Pathway level impacts of in situ exposure of fathead minnow to effluent mixtures in the Duluth harbor

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There is a major effort to characterize the potential adverse effects of effluents released from sewage treatment plants in North America. At many locations pharmaceuticals, endocrine disruptors, and other chemicals of emerging concerns are present in the environment at concentrations capable of altering physiological processes in wildlife. In order to elucidate the effects of sewage treatment plants on wildlife, fathead minnows were caged in situ at several locations at the Duluth (MN) harbor, proximal and distal from the wastewater treatment plant discharge location. Fish were also exposed in the laboratory to 100% of effluent as a positive control. Several endpoints such as steroid hormones, vitellogenin, transcriptomics and metabolomics were analyzed. We applied a pathway level approach to understand the effects of the discharge as well as the potential dilution of the adverse effects at distal locations. Effects on gene expression were very similar for the different locations, with a higher amount of changes for the 100% effluent and the higher intensity for fish exposed to locations closer to the discharge site. Changes in gene expression were involved in pathways related to cytochrome P450s, hypoxia inducible factor, neurological function, or embryonic development among others. While distance from the discharge site decreased the effects on gene expression, pathway analysis was very effective in detecting the potential adverse effects of the discharge.

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STICs Field	Entry		
1 – Influence/profile	Not applicable		
2 – Clearance tracking	Assigned automatically		
no.			
3 – Principal Investigator	Gerald Ankley		
/ Project Officer			
4- Product title	Copy and paste from abstract		
5 - Authors	See abstract		
6a- Product type	Presentations and technical summaries		
6b-Product subtype	Abstract		
6c – Records schedule	Not a senior official		
7a – Impact statement	n/a		
7b- Product description	Paste in abstract		
8 – Bibliographic citation	SETAC North America 33rd Annual Meeting, 11-15 November,		
0. 4	Long Beach, CA, USA. Public		
9 - Access	2.1.2 2.1.2: AOP-based effects monitoring and exposure reconstruction		
10 – Tracking and	2.1.2 2.1.2. Not based chools mornioning and exposure reconstitution		
Planning Task			
10 – Tracking and	(2) Case studies evaluating the utility of transcriptomics, metabolomics, and		
Planning	associated bioinformatic methods for comparing the nature and severity of		
Product	biological impairment as a function of space and/or time to assess the		
	efficacy of remediation efforts within Great Lakes Areas of Concern.		
11 – Copyright	No		
permission			
12 - QA	not applicable		
13 – Policy implications	No		
14 - Keywords	transcriptomics		
	effects-based monitoring		
	wastewater		
	mixture		
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